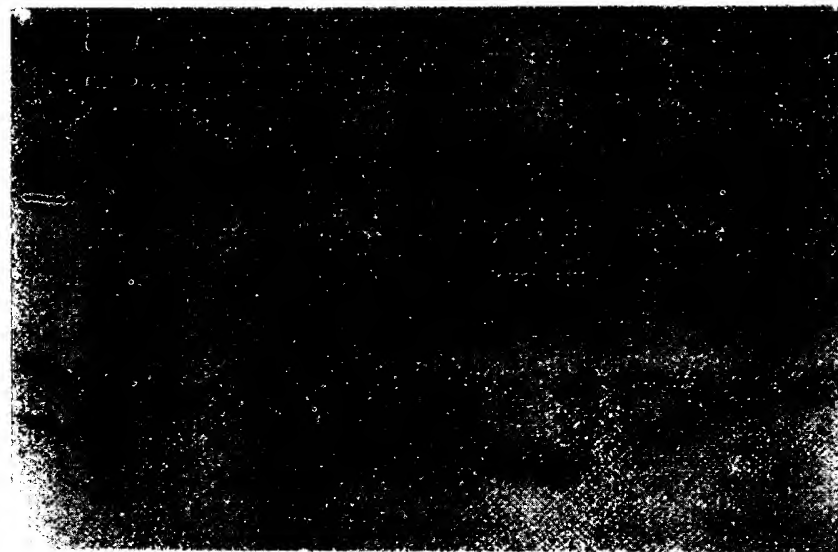


fraction: M L 21 22 23 24 25 26 27 28 29 30 31 32

70 kD

30 kD

18 kD



SCG SURVIVAL

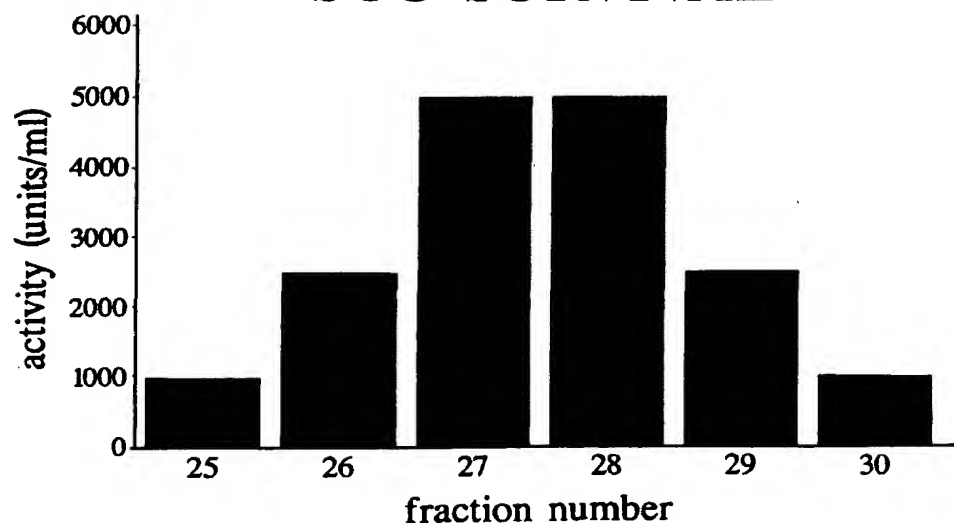
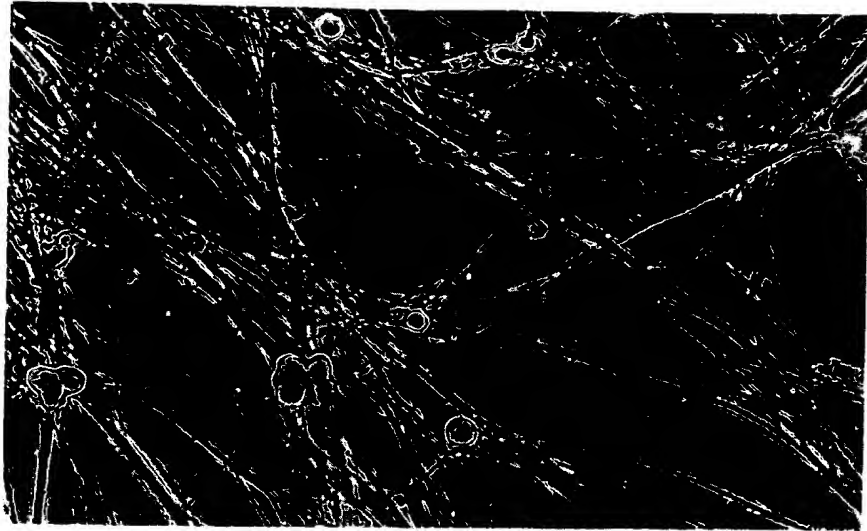
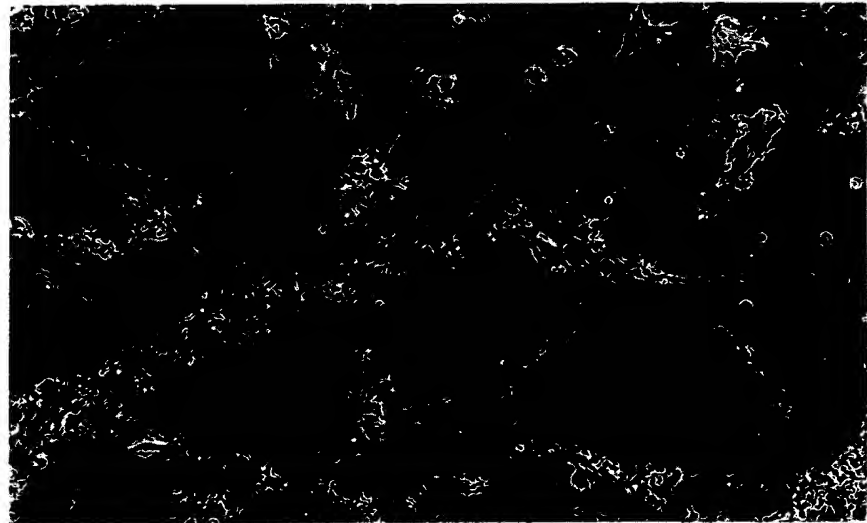


Figure 2

A) NGF



B) Anti-NGF



C) Anti-NGF
+
Neurturin

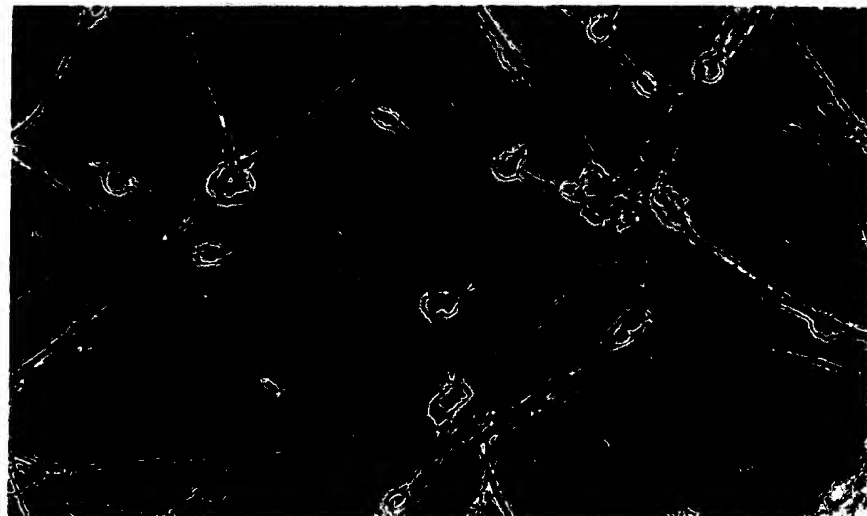


Figure 3

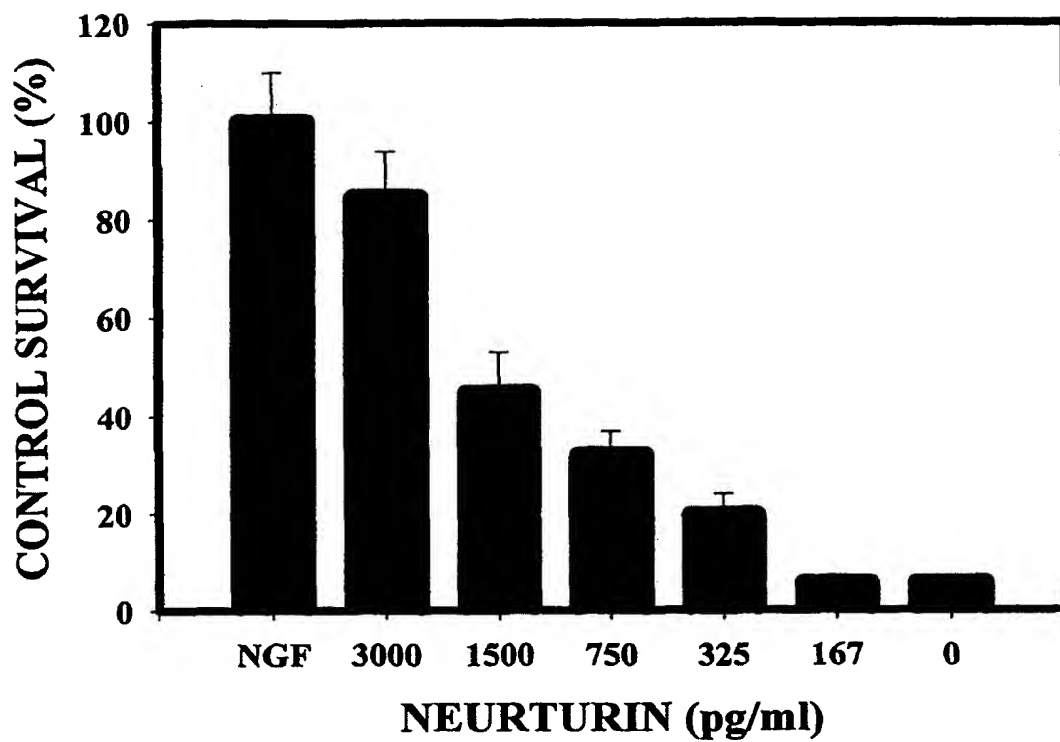


Figure 4

1	S P O K Q M A V L P R R E R N R Q A A A A N P E N S R G K G	hGDNF
1	S P O K Q A A A L P R R E R N R Q A A A A S P E N S R G K G	mGDNF
1	S P O K Q A A A L P R R E R N R Q A A A A S P E N S R G K G	rGDNF
1	- -	hNTN
1	- -	mNTN
31	R R G Q R G K N R G C V L T A I H L N V T D L G L G Y E T K	hGDNF
31	R R G Q R G K N R G C V L T A I H L N V T D L G L G Y E T K	mGDNF
31	R R G Q R G K N R G C V L T A I H L N V T D L G L G Y E T K	rGDNF
1	- - - A R L G A R P C G L R E L E V R V S E L G L G Y A S D	hNTN
1	- - - - - P G A R P C G L R E L E V R V S E L G L G Y T S D	mNTN
61	E E L I F R Y C S G S C D A A E T T Y D K I L K N L S R N R	hGDNF
61	E E L I F R Y C S G S C E A A E T M Y D K I L K N L S R S R	mGDNF
61	E E L I F R Y C S G S C E A A E T M Y D K I L K N L S R S R	rGDNF
28	E T V L F R Y C A G A C E A A A R V Y D L G L R R L R Q R R	hNTN
26	E T V L F R Y C A G A C E A A A I R I Y D L G L R R L R Q R R	mNTN
91	R L V S O K V - G Q A C C R P I A F D D D L S F L D D N L V	hGDNF
91	R L T S O K V - G Q A C C R P V A F D D D L S F L D D N L V	mGDNF
91	R L T S O K V - G Q A C C R P V A F D D D L S F L D D S L V	rGDNF
58	R L R R E R V R A Q P C C R P T A Y E D E V S F L D A H S R	hNTN
56	R V R R E R A R A H P C C R P T A Y E D E V S F L D V H S R	mNTN
120	Y H I L R K H S A K R C G C I .	hGDNF
120	Y H I L R K H S A K R C G C I .	mGDNF
120	Y H I L R K H S A K R C G C I .	rGDNF
88	Y H T V H E L S A R E C A C V .	hNTN
86	Y H T L Q E L S A R E C A C V .	mNTN

Figure 5

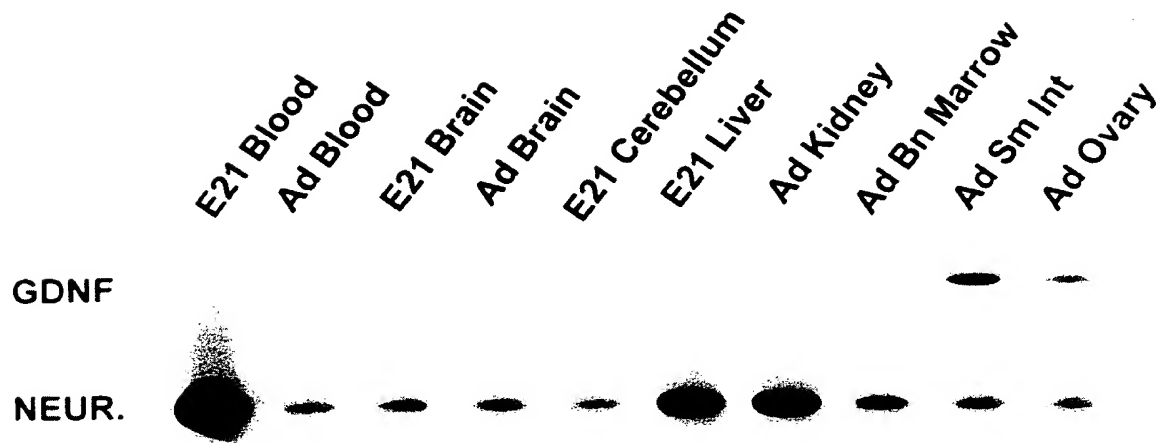


Figure 6

ATGCAGCGCTGGAAGGCGGCGGCCTTGGCCTCAGTGCTCTGCAGCTCCGTGCTGTCCATC 60
Met Gln Arg Trp Lys Ala Ala Ala Leu Ala Ser Val Leu Cys Ser Ser Val Leu Ser Ile

TGGATGTGTCGAGAGGGCCTGCTTCTCAGCCACCGCCTCGGACCTGCGCTGGTCCCCCTG 120
Trp Met Cys Arg Glu Gly Leu Leu Leu Ser His Arg Leu Gly Pro Ala Leu Val Pro Leu

CACCGCCTGCCTCGAACCCTGGACGCCCCGATTGCCCGCCTGGCCCAGTACCCTGCACTC 180
His Arg Leu Pro Arg Thr Leu Asp Ala Arg Ile Ala Arg Leu Ala Gln Tyr Arg Ala Leu

CTGCAGGGGGCCCCGGATGCGATGGAGCTGCGCGAGCTGACGCCCTGGGCTGGGCGGCCCC 240
Leu Gln Gly Ala Pro Asp Ala Met Glu Leu Arg Glu Leu Thr Pro Trp Ala Gly Arg Pro

CCAGGTCCGCGCCGTGCGGCGGGGCCCCGGCGGGCGCGCGCGTGC GCGGGTTGGGGGCG 300
Pro Gly Pro Arg Arg Arg Ala Gly Pro Arg Arg Arg Arg Ala Arg Ala Arg Leu Gly Ala

CGGCCTTGCGGGCTGCGCGAGCTGGAGGTGCGCGTGAGCGAGCTGGGCCTGGGCTACGCG 360
Arg Pro Cys Gly Leu Arg Glu Leu Glu Val Arg Val Ser Glu Leu Gly Leu Gly Tyr Ala

TCCGACGAGACGGTGCTGTTCGCTACTGCGCAGGCGCCTGCGAGGCTGCCGCGCGCGTC 420
Ser Asp Glu Thr Val Leu Phe Arg Tyr Cys Ala Gly Ala Cys Glu Ala Ala Ala Arg Val

TACGACCTCGGGCTGCGACGACTGCGCCAGCGGCGGCGCCTGCGGCGGGAGCGGGTGCGC 480
Tyr Asp Leu Gly Leu Arg Arg Leu Arg Gln Arg Arg Arg Leu Arg Arg Glu Arg Val Arg

GCGCAGCCCTGCTGCGGCCCGACGCGCTACGAGGACGAGGTGTCCTTCTGGACGCGCAC 540
Ala Gln Pro Cys Cys Arg Pro Thr Ala Tyr Glu Asp Glu Val Ser Phe Leu Asp Ala His

AGCCGCTACCACACGGTGACGAGCTGTCGGCGCGCGAGTGCGCCTGCGTGTGA 594
Ser Arg Tyr His Thr Val His Glu Leu Ser Ala Arg Glu Cys Ala Cys Val

Figure 7

ATGAGGCGCTGGAAGGCAGCGGCCCTGGTGTGCTCATCTGCAGCTCCCTGCTATCTGTC 60
Met Arg Arg Trp Lys Ala Ala Ala Leu Val Ser Leu Ile Cys Ser Ser Leu Leu Ser Val

TGGATGTGCCAGGAGGGTCTGCTCTTGGGCCACCGCCTGGGACCCGCGCTTGCCCCGCTA 120
Trp Met Cys Gln Glu Gly Leu Leu Leu Gly His Arg Leu Gly Pro Ala Leu Ala Pro Leu

CGACGCCCTCCACGCACCCTGGACGCCCGCATCGCCCGCCTGGCCCA[↓]GTATCGCGCTCTG 180
Arg Arg Pro Pro Arg Thr Leu Asp Ala Arg Ile Ala Arg Leu Ala Gln Tyr Arg Ala Leu

CTCCAGGGCGCCCCGACGCGGTGGAGCTTCGAGAACTTCTCCCTGGGCTGCCCGCATC 240
Leu Gln Gly Ala Pro Asp Ala Val Glu Leu Arg Glu Leu Ser Pro Trp Ala Ala Arg Ile

CCGGGACCGCGCCGTCGAGCGGGTCCCCGGCGTCGGCGGGCGCGGGCCGGGGGCTCGGCCT 300
Pro Gly Pro Arg Arg Arg Ala Gly Pro Arg Arg Arg Arg Ala Arg Pro Gly Ala Arg Pro

TGTGGGCTGCGCGAGCTCGAGGTGCGCGTGAGCGAGCTGGGCCTGGGCTACACGTCTGGAT 360
Cys Gly Leu Arg Glu Leu Glu Val Arg Val Ser Glu Leu Gly Leu Gly Tyr Thr Ser Asp

GAGACCGTGCTGTTCCGCTACTGCGCAGGCGCGTGCGAGGCGGCCATCCGCATCTACGAC 420
Glu Thr Val Leu Phe Arg Tyr Cys Ala Gly Ala Cys Glu Ala Ala Ile Arg Ile Tyr Asp

CTGGGCCTTCGGCGCCTGCGCCAGCGGAGGCGCGTGCGCAGAGAGCGGGCGCGGGCGCAC 480
Leu Gly Leu Arg Arg Leu Arg Gln Arg Arg Arg Val Arg Arg Glu Arg Ala Arg Ala His

CCGTGTTGTGCCCCGACGGCCTATGAGGACGAGGTGTCCTTCTGGACGTGCACAGCCGC 540
Pro Cys Cys Arg Pro Thr Ala Tyr Glu Asp Glu Val Ser Phe Leu Asp Val His Ser Arg

TACCACACGCTGCAAGAGCTGTCGGCGCGGGAGTGCGCGTGCGTGTGA 588
Tyr His Thr Leu Gln Glu Leu Ser Ala Arg Glu Cys Ala Cys Val

Figure 8

GGAGGGAGAGCGCGCGGTGGTTTCGTCCGTGTGCCCCGCGCCCGGCGC	-301
TOCTCGCGTGGCCCCGCGTCCTGAGCGCGCTCCAGCCTCCACGCGCGCC	-251
ACCCCGGGGTTCACTGAGCCCGGCGAGCCCGGGGAAGACAGAGAAAGAGA	-201
GGCCAGGGGGGGAACCCCATGGCCCGGCCCGTGTCCCGCACCCCTGTGCGG	-151
TGGCCTCCTCCGGCACGGGGTCCCCGGGTGCGCTCCGGTCCCCGCGATCC	-101
GGATGGCGCACGCAGTGGCTGGGGCCGGGCCGGGCTCGGGTGGTGGGAGG	-51
AGTCACCACTGACCGGGTCATCTGGAGCCCGTGGCAGGCCGAGGCCAGG	-1
<u>ATGAGGCGCTGGAAGGCAGCGGCCCTGGTGTGCTCATCTGCAGCTCCCT</u>	50
<u>GCTATCTGTCTGGATGTGCCAGGAGGGTCTGCTCTTGGGCCACCGCCTGG</u>	100
<u>GACCCGCGCTTGCCCCGCTACGACGCCCTCCACGCACCCCTGGACGCCCGC</u>	150
<u>ATCGCCCGCCTGGCCAGTATCGCGCTCTGCTCCAGGGCGCCCCGACGC</u>	200
<u>GGTGGAGCTTCGAGAACTTTCTCCCTGGGCTGCCCGCATCCCGGGACCGC</u>	250
<u>GCCGTGAGCGGGTCCCCGGCGTGGCGGGCGCGGCCGGGGGCTCGGCCT</u>	300
<u>TGTGGGCTGCGCGAGCTCGAGGTGCGCGTGAGCGAGCTGGGCCTGGGCTA</u>	350
<u>CAGTCCGATGAGACCGTGCTGTTCCGCTACTGCGCAGGCGCGTGCGAGG</u>	400
<u>CGGCCATCCGCATCTACGACCTGGGCCTTCGGCGCCTGCGCCAGCGGAGG</u>	450
<u>CGCGTGCGCAGAGAGCGGGCGCGGGCGCACCCGTGTTGTGCCCCGACGGC</u>	500
<u>CTATGAGGACGAGGTGTCTTCTGACGTGCACAGCCGCTACCAACACGC</u>	550
<u>TGCAAGAGCTGTGCGCGCGGGAGTGCGCGTGCGTGTGATGCTACCTCACG</u>	600
CCCCCGACCTGCGAAAGGGCCCTCCCTGCCGACCCTCGCTGAGAACTGA	650
CTTCACATAAAGTGTGGGAACTCCC	675

Figure 9

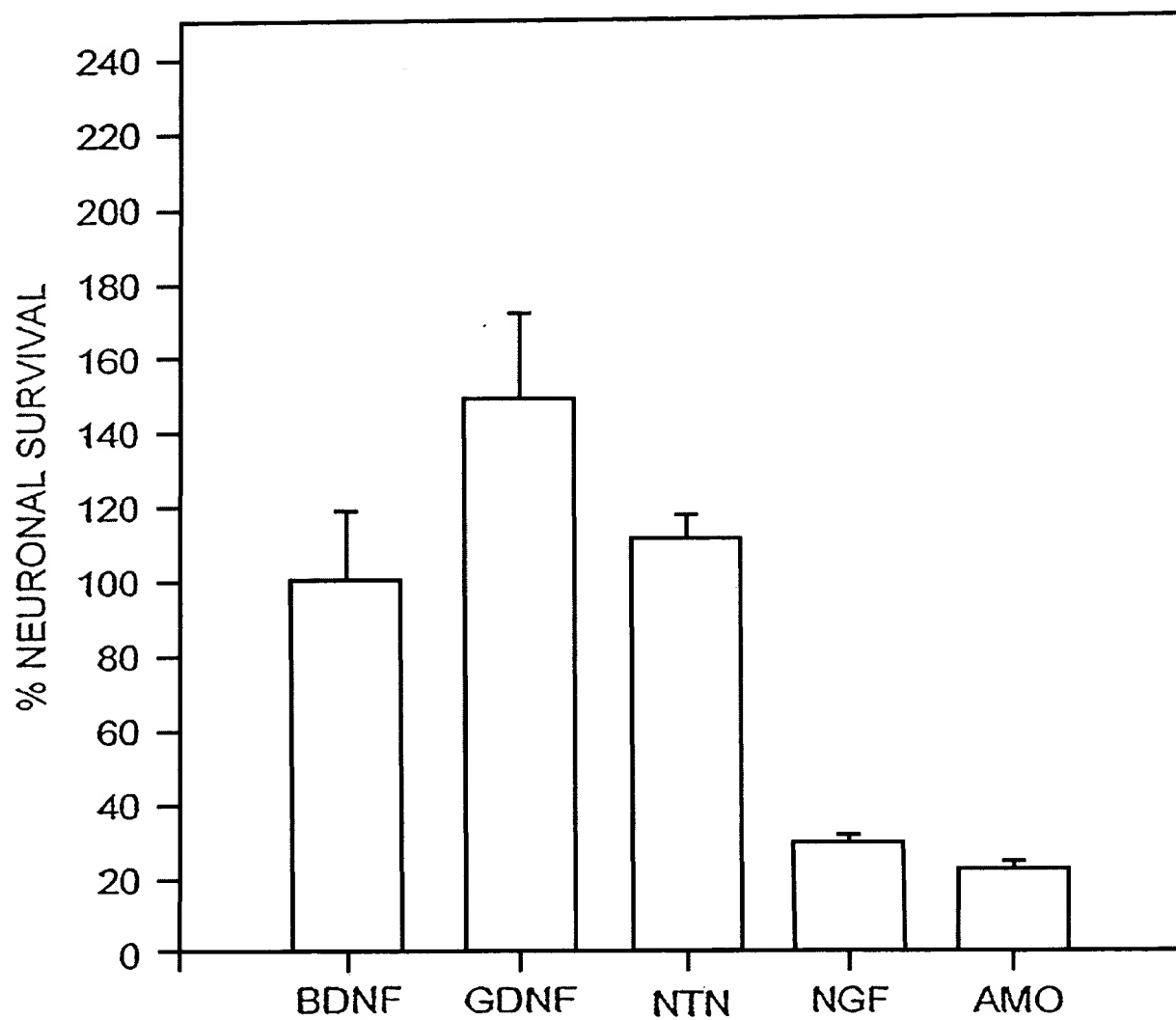


Figure 10

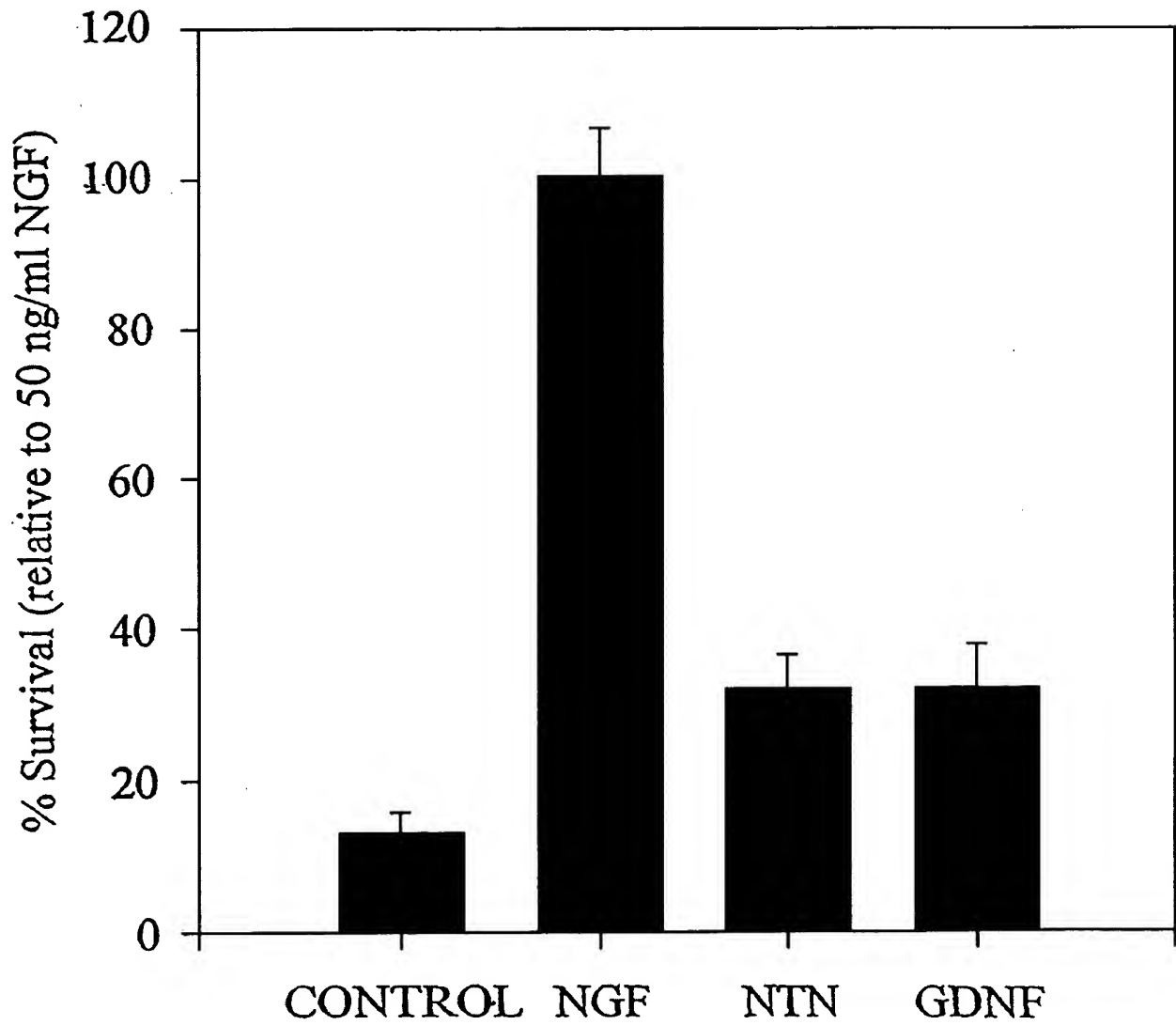


Figure 11

Title:
Inventor(s):
Appln. No.
Docket #

Neurturin and Related Growth Factors
Johnson et al.
09/476,290 - Replacement Sheet
56029-2669

12 / 19

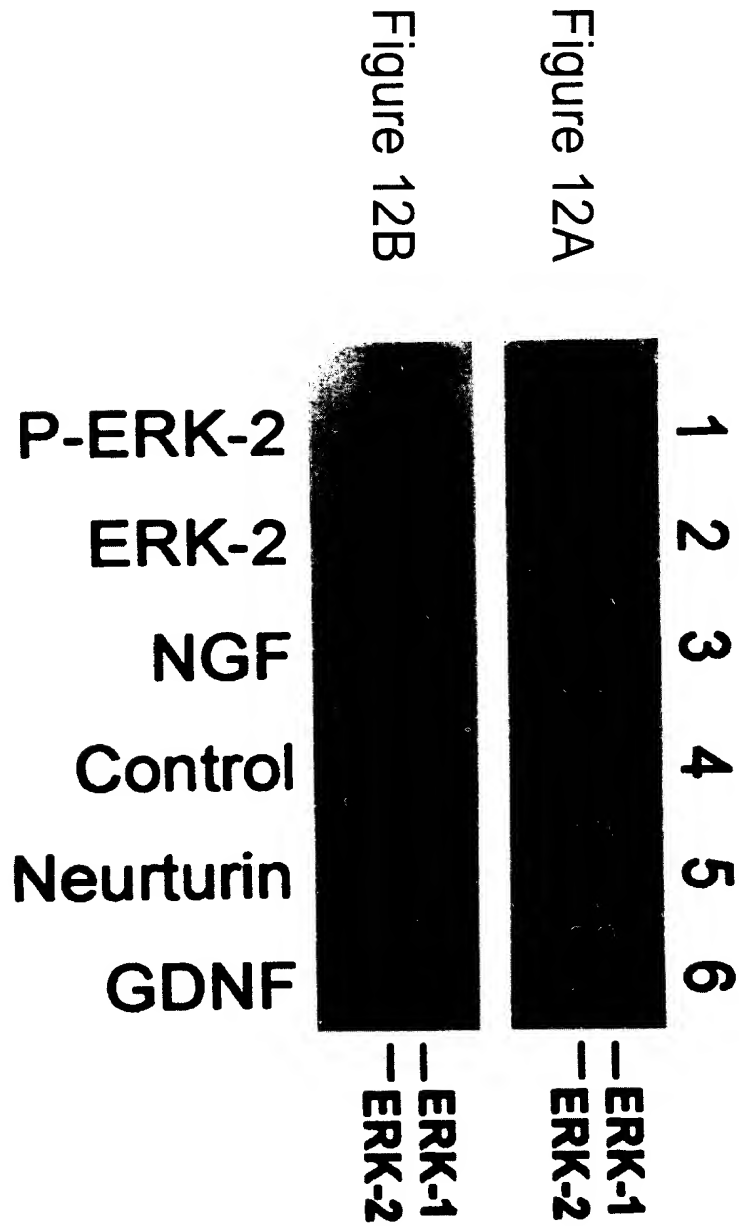
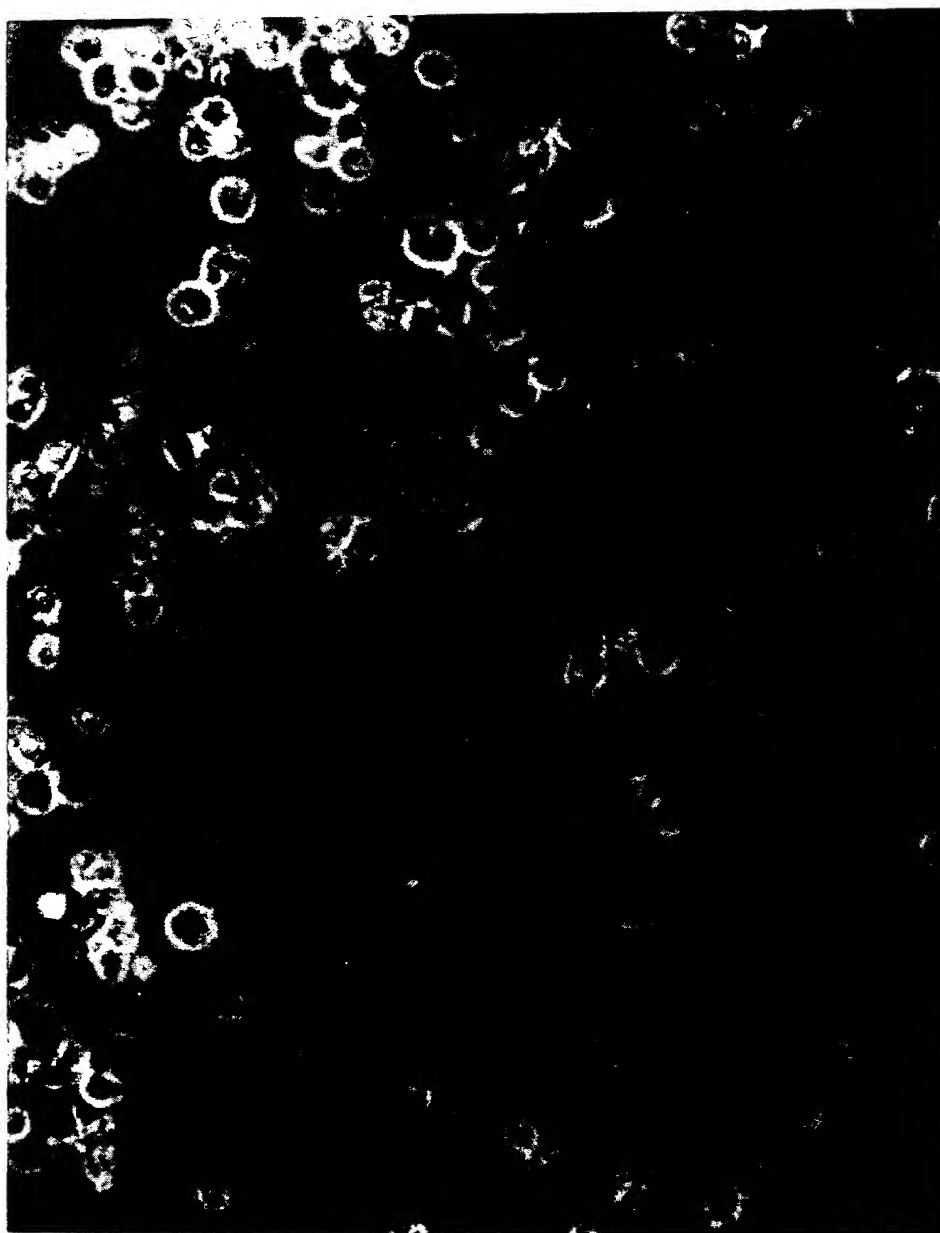


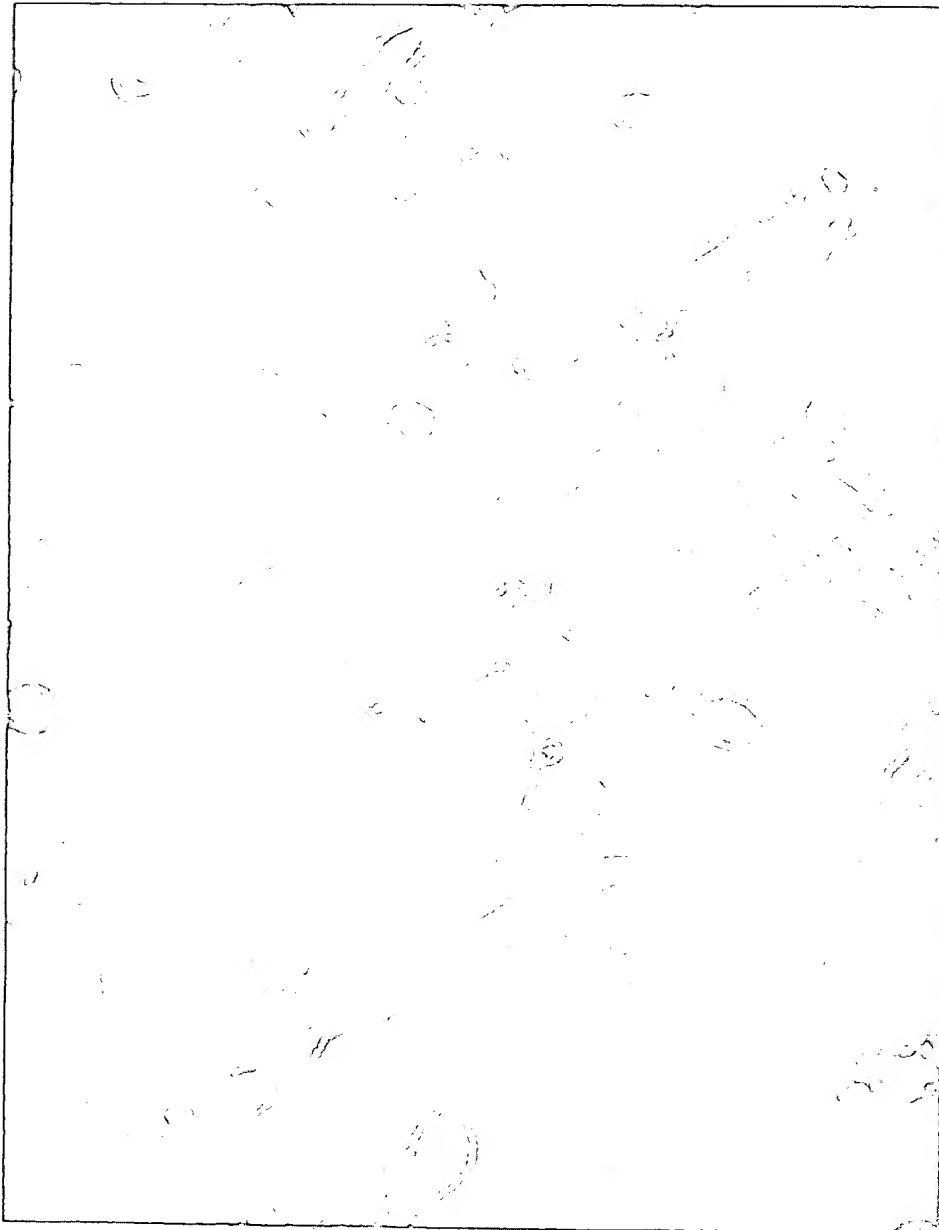
Figure 13A. Untreated



Title: Neurturin and Related Growth Factors
Inventor(s): Johnson et al.
Appl. No. 09/476,290 - Replacement Sheet
Docket # 56029-2669

14 / 19

Figure 13B. Neurturin-treated



MAPK Response in Neuroblastoma Cell Lines

Figure 14A

SK-NSH Neuroblastoma (naive)

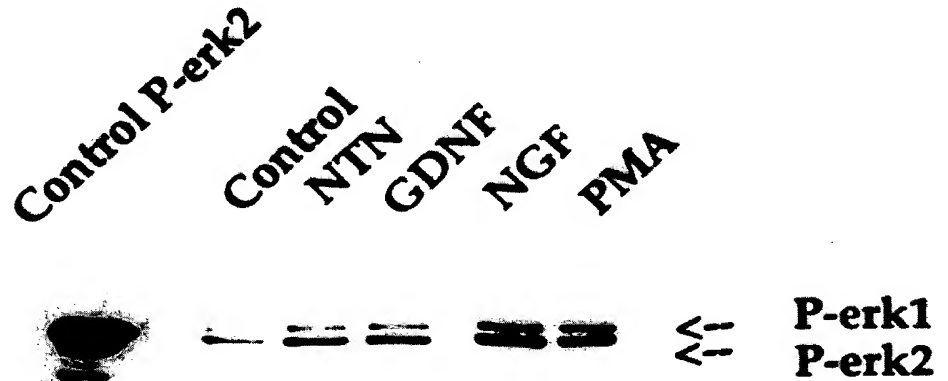


Figure 14A

Figure 14B **NGP Neuroblastoma (RA tx)**

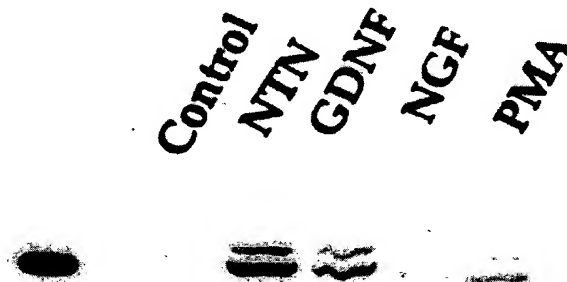
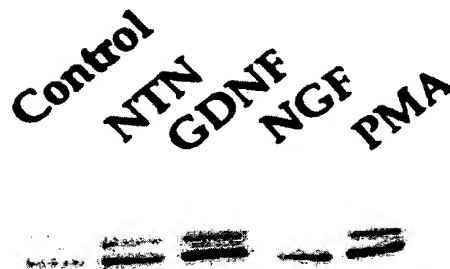


Figure 14C **SY5Y Neuroblastoma (RX tx)**



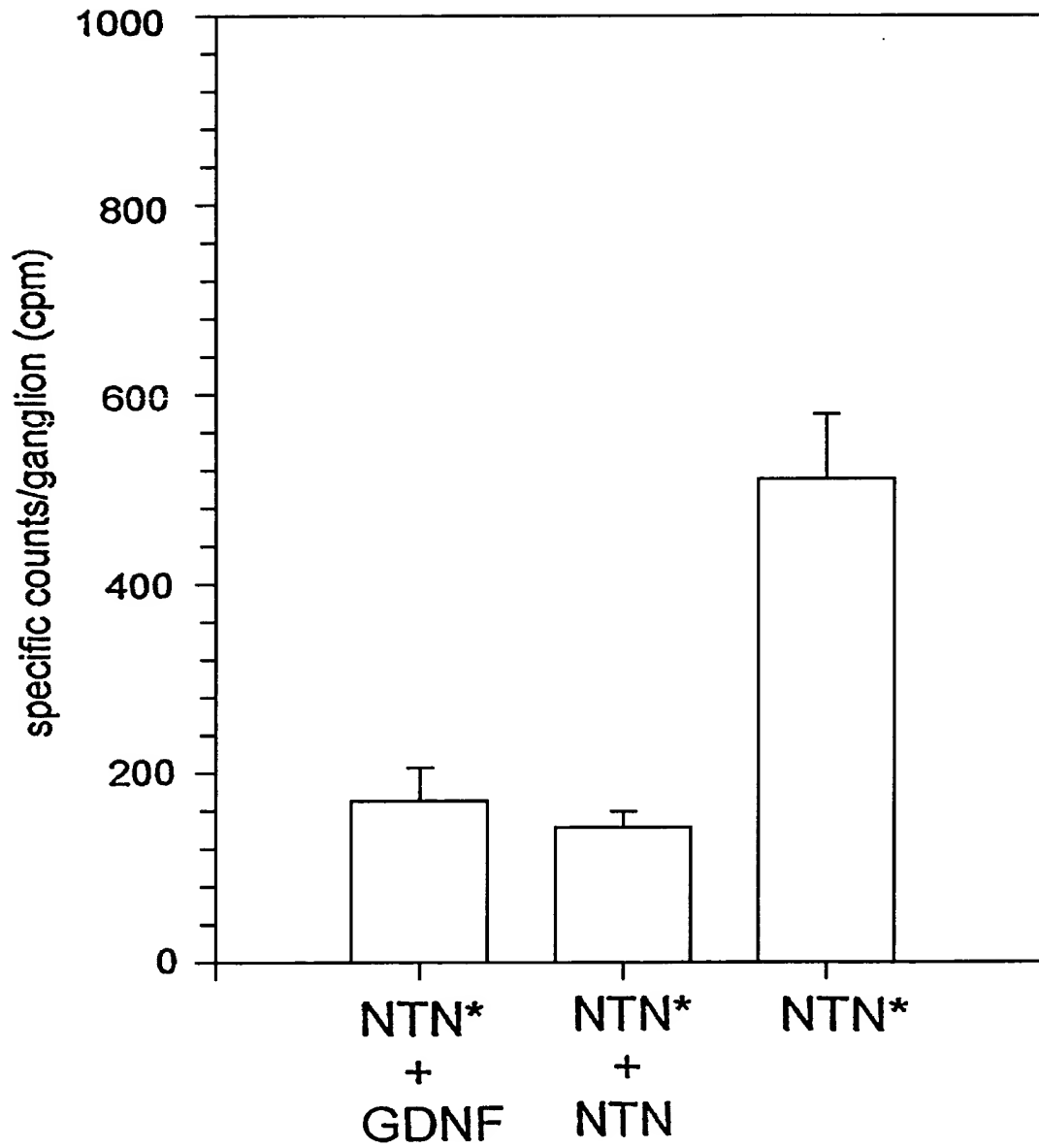


Figure 15

SEQ ID NO:	GROWTH FACTOR	SEQUENCE
134	TGFβ1	CCVRQLYIDFRKDLGWR-WIHEPKGYANFCLGRCPIYNSLDT-----QYSKVLALYNQENPGASAA-PCCV--PQALEPLPIVYVYGRKPKV--EQLSNMIVRSCKCS
135	TGFβ2	CCIRPLYIDFRKDLGWR-WIHEPKGYANFCAACAPYIWSSDT-----QHSKVLALYNQENPGASAA-PCCV--SODLEPLTILYVYIGKPKI--EQLSNMIVRSCKCS
136	TGFβ3	CCVRPLTYIDFRQDLGWR-WIHEPKGYANFCAACAPYIWSSDT-----THSTVLGLYNTINPEASAS-PCCV--PQLEPLTILYVYIGKPKV--EQLSNMIVRSCKCS
137	INHβA	CCCKQFVSFK-DIGWMDMIIAPSGYANFCEGRCPSHLAG-TSGSGLS9THSVTINHYRMGHH3PANKSCCV--PTLRPMNMLYDDGQNTI-KKDIONMIVEECGCS
138	INHβB	CCRCQFFIDFR-LIGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
139	NODAL	CRVKPQVDFN-LIGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
140	BMP2	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
141	BMP4	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
142	DDP	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
143	BMP5	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
144	BMP6	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
145	BMP7	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
146	BMP8	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
147	60A	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
148	BMP3	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
149	VG1	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
150	GDF1	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
151	GDF3	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
152	DORSALN	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
153	INHα	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
154	MIS	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
155	GDF9	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
156	GDNF	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA
157	NTN	CRHSLYVDPS-DVGWMDMIIAPSGYANFCEGRCPSHLAG-VPQSAS9FTEAVNQRMGLTF-GTVNSCCI--PTLSTMSMLYFDDENIV-KRDVPMNIVEECGCA

Figure 16

SEQ ID NO:	GROWTH FACTOR	SEQUENCE	
86	TGFβ1	CCVRQLYIDFRKDLGWK-WIHEPKGYHANFCLGPCPYIWSLDT-----QYSKVLALYNQHNPGASAA-P	62
87	TGFβ2	CCLRPLYIDFKRDLGWK-WIHEPKGYNANFCAGACPYLMSBDT-----QHSRVLSLYNTINPEASAS-P	62
88	TGFβ3	CCVRPLYIDFRDLGWK-WIHEPKGYNANFCAGACPYLMSBDT-----THSTVLGLYNTLNPEASAS-P	62
89	INHα	CKKQFFVSPK-DIGWMDWIAPSGYHANYCEGECPSHIAJ-TSGSLSFHSIVINHRYMRGHSPPANLKS	69
90	INHβ	CCRQFPIDFR-LIGWMDWIAPTYGNYCEGSCPAYLAS-VPGBASBHTAVNQYRMQLNP-GTVNS	68
91	NODAL	CRVKFQVDFN-LIGWMDWIAPKQYNAYRCBGCPCNPVGESEFHT-----NHAYIQSLKRYQPHR-VPST	65
92	BMP2	CKRHPLYVDFS-DVGWMDWIVAPPGYHAFYCHGECPPPLADHLNST-----NHAIVQTLVNSVNS-K-IPKA	64
93	BMP4	CRHSLYVDFS-DVGWMDWIVAPPGYHAFYCHGDCPPPLADHLNST-----NHAIVQTLVNSVNS-S-IPKA	64
94	DPP	CRHSLYVDFS-DVGWMDWIVAPPGYHAFYCHGKCPPLADHLNST-----NHAVQTLVNNANPGK-VPKA	65
95	BMP5	CKKHELVSFR-DIGWMDWIAPEGYAAFYCDGECSPPLAHMNAT-----NHAIVQTLVHLMPDH-VPKP	65
96	BMP6	CKKHELVSFR-DIGWMDWIAPKGYAANYCDGECSPPLAHMNAT-----NHAIVQTLVHLMPDH-VPKP	65
97	BMP7	CKKHELVSFR-DIGWMDWIAPEGYAAFYCEGECAPPLNSYNAT-----NHAIVQTLVHLMPDH-VPKP	65
98	BMP8	CRHSLYVDFQ-DIGWMDWIVAPPGYSAFYCEGECSPPLNSYNAT-----NHAIQSLVHLMPDH-VPKA	65
99	60A	COMQTLYIDFK-DIGWMDWIAPEGYAAFYCEGECSPPLNSYNAT-----NHAIQSLVHLMPDH-VPKA	65
100	BMP3	CARLYLKVDFQ-DIGWMDWIAPKGYAANYCDGECSPPLAHMNAT-----NHAIQSLVHLMPDH-VPKA	65
101	VG1	CKKRLYVDFK-DVGWMDWIVAPPGYHAFYCHGECPPPLADHLNST-----NHAIQSLVHLMPDH-VPKA	65
102	GDF1	CRARLYVDFR-EVGWMDWIVAPPGYHAFYCHGECPPPLADHLNST-----NHAIQSLVHLMPDH-VPKA	69
103	GDF3	CHRHQLFINFQ-DIGWMDWIVAPPGYHAFYCHGECPPPLADHLNST-----NHAIQSLVHLMPDH-VPKA	64
104	DORSAL	CRRTSLHVNFK-EIGWMDWIVAPPGYHAFYCHGECPPPLADHLNST-----KHAIQSLVHLMPDH-VPKA	65
105	INHα	CHRVALNISF-QELGMDWIVAPPGYHAFYCHGECPPPLADHLNST-----KHAIQSLVHLMPDH-VPKA	65
106	MIS	CALRELSVDLRAERS---VLIPTYQANNCQAGCWPQSDR---NPRYGNHVVILLKMOARGATLARP	63
107	GDF9	CELHDFSLFS-QLKMDWIVAPPGYHAFYCHGECPPPLADHLNST-----HTVQNMITYE-KLDPVSPSP	65
108	GDNF	CVLTAHLNVT-DLGLG--YETKEELIPRYCSGSCD-AAETTYDKILKNLSRN-----RRLVSDKV-GQA	60
109	NTN	CGRLRELVRS-ELGLG--YASDETVLFRYCAGACE-AAARVYDLGLRLRQR-----RRLRBRVRARQP	61

Figure 17

SEQ ID NO:	GROWTH FACTOR	SEQUENCE
110	TGF β 1	CCV--PQALEPLPIVYYVGRKPKV--EQLSNMIVRSCKCS
111	TGF β 2	CCV--SQDLEPLTILYYIGKTPKI--EQLSNMIVKSCKCS
112	TGF β 3	CCV--PQDLEPLTILYYVGRTPKV--EQLSNMVVKSCCKCS
113	INH β A	CCV--PTKLRPMSMLYYDDGQNI I-KKDIQNMIVEECGCS
114	INH β B	CCI--PTKLSTMSMLYFDDEYNIV-KRDVPNMIVEECGCA
115	NODAL	CCA--PVKTKPLSMLYVDNGR--VLLEHHKDMIVEECGCL
116	BMP2	CCV--PTELSAISMLYLDENEKVVVK-NYQDMVVEGCGCR
117	BMP4	CCV--PTELSAISMLYLDDEYDKVVVK-NYQEMVVEGCGCR
118	DPP	CCV--PTQLDSVAMLYLNDQSTVVVK-NYQEMTVVGCGCR
119	BMP5	CCA--PTKLNAISVLYFDDSSNVILK-KYRNMVVRSCGCH
120	BMP6	CCA--PTKLNAISVLYFDDNSNVILK-KYRNMVVRACGCH
121	BMP7	CCA--PTQLNAISVLYFDDSSNVILK-KYRNMVVRACGCH
122	BMP8	CCA--PTKLSATSVLYYDSSNNVILR-KHRNMVVKACGCH
123	60A	CCA--PTRLGALPVLYHLNDENVNLK-KYRNMIVKSCGCH
124	BMP3	CCV--PEKMSSLSILFFDENKNVVVKV-YPNMTVESACR
125	VG1	CCV--PTKMSPISMLFYDNNDNVVLR-HYENMAVDECGCR
126	GDF1	CCV--PARLSPISVLFFDNSDNVVLR-QYEDMVVDECGCR
127	GDF3	VCV--PTKLSPISMLYQDSKKNVILR-HYEDMVVDECGCG
128	DORSLN	CCV--PTKLDAISILYKDDAGVPTLIYNYEGMKVAECGCR
129	INH α	CCAALPGTMRPLHVRTTSDGGYSFKYETVPNLLTQHCACI
130	MIS	CCV--PTAYT--GKLLISLSEERISAHVPMNVATECGCR
131	GDF9	SCV--PGKYSPLSVLTIEPDGSIAYK-EYEDMMATSCTCR
132	GDNF	CCRPIAFD-DDLSFL-----DDNLVYHILRKHSACRCGCI
133	NTN	CCRPTAYE-DEVSFL-----DAHSRYHTVHEL SARECACV

Figure 18

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